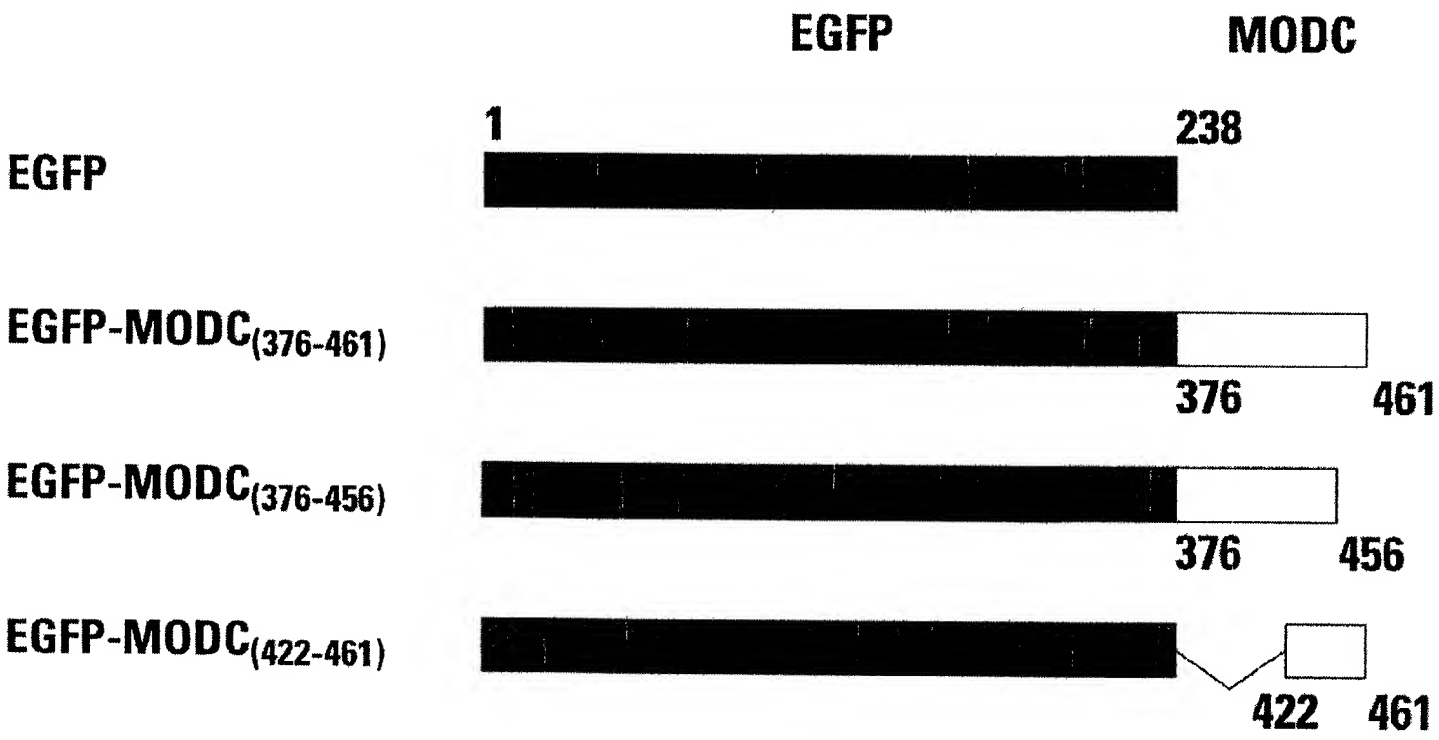

FIG. 1

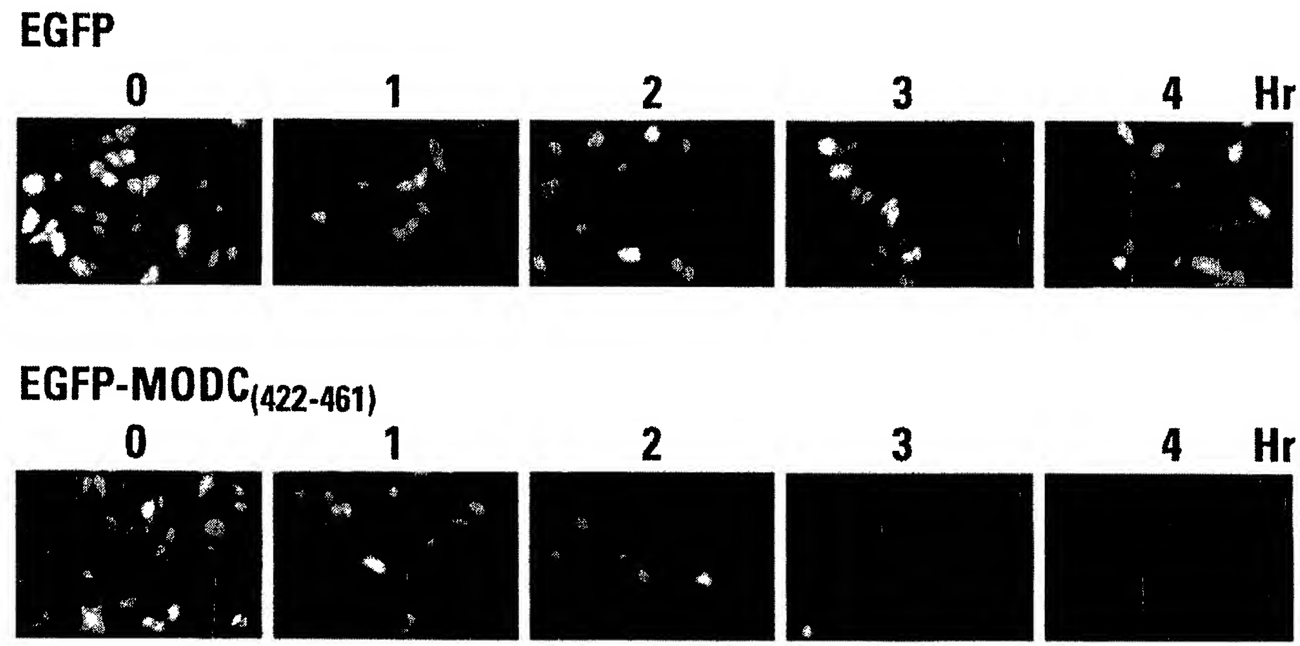

FIG. 2

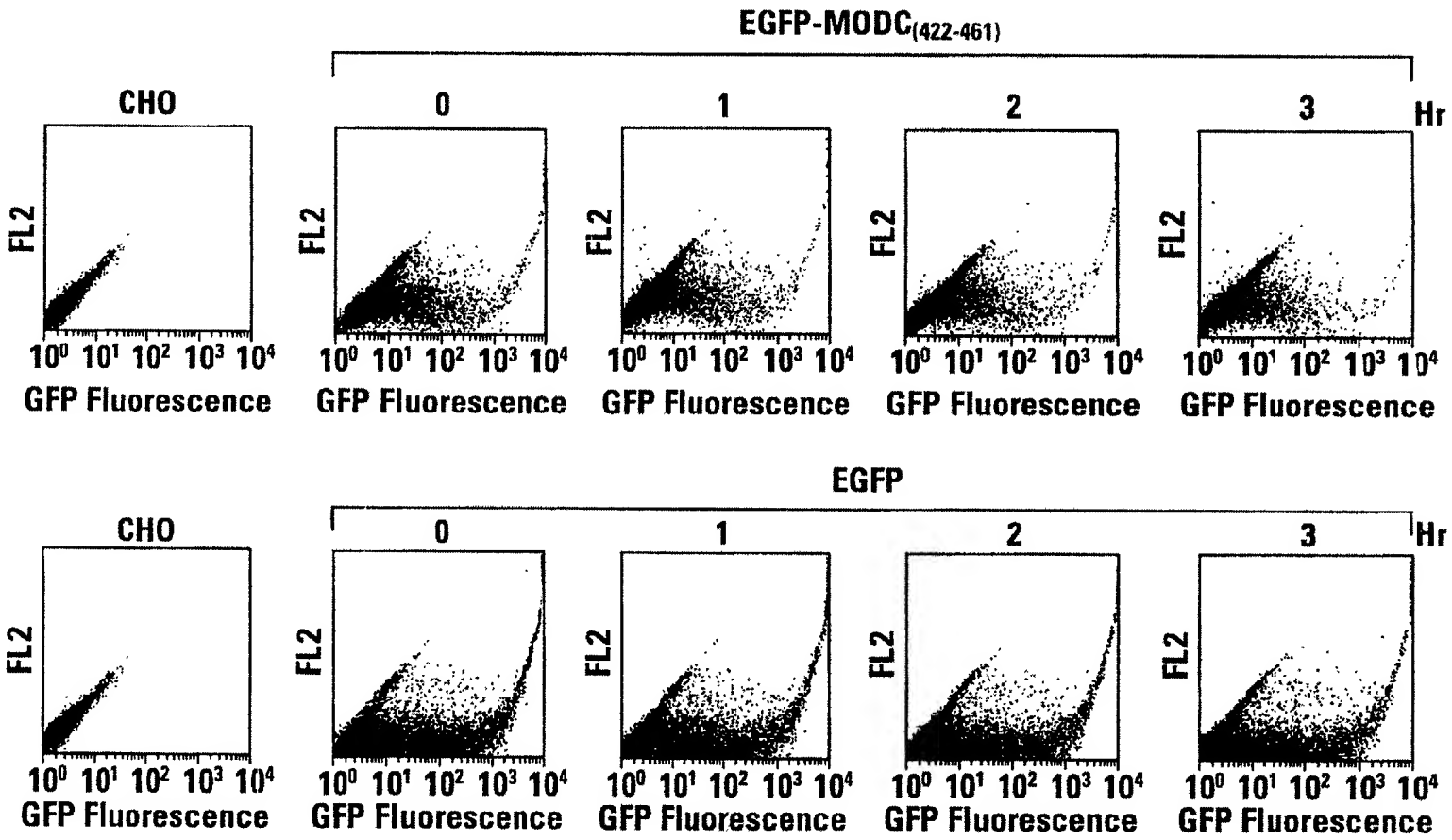

FIG. 3

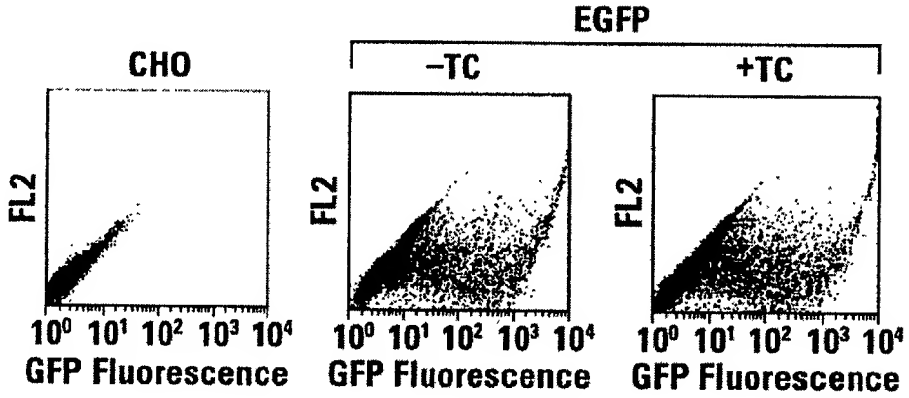

FIG. 3 cont.

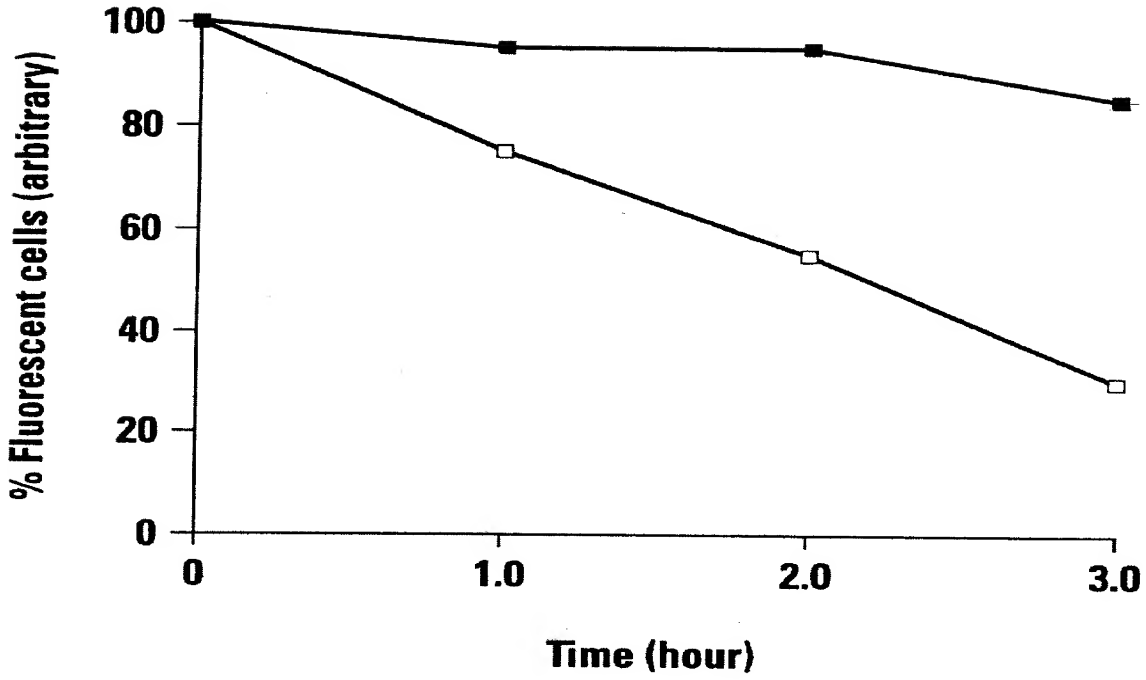

FIG. 4

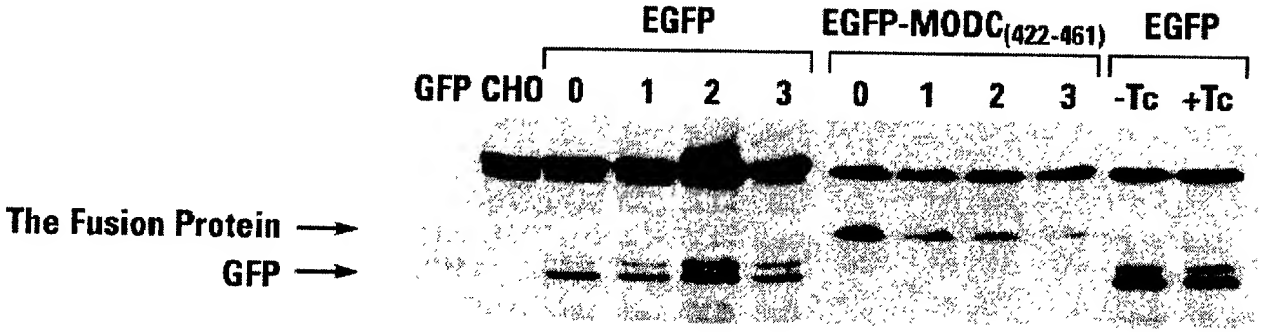

FIG. 5

PEST Sequence

	423	450	
EGFP-MODC$_{422-461}$	--HGF**PP**E**V**E**E**Q**DD**G**T**L**P**M**S**CAQ**ES**GM**DRH**--		(SEQ ID NO. 3)
P426A/P427A	-----**AA**----------		
P438A	----------**A**----------		
E428A/E430A/E431A	-------**A**-**AA**----------		
E44A	----------**A**--------		
S440A	----------**A**----------		
S445A	----------**A**-------		
T436A	----------**A**----------		
D433A/D434A	----------**AA**----------		
D448A	----------**A**----		
H423A	--**A**----------		
R449A/H450A	----------**AA**--		

FIG. 6

FIG. 7

Table 1. FACS analysis of EGFP, EGFP-MODC $_{422-461}$, and mutations in transfected CHO K1 Tet-off cells.

Constructs	0h	(initial)	2h	4h
EGFP	100%	(63.6)	107%	92%
EGFP-MODC $_{422-461}$	100%	(12.6)	52%	29%
P426A/P427A	100%	(11.5)	39%	11%
P438A	100%	(34.1)	79%	60%
E428A/E430A/E431A	100%	(17.3)	20%	15%
E444A	100%	(12.6)	69%	65%
S440A	100%	(21.6)	78%	66%
S445A	100%	(23.5)	29%	20%
T436A	100%	(46.9)	70%	47%
D433A/D434A	100%	(11.31)	22%	6%
D448A	100%	(32.6)	30%	15%
H423A	100%	(12.2)	50%	25%
R449A/H450A	100%	(27.9)	93%	86%

Transfection was performed in CHO/tTA cells using the procedure as described in Methods. After 24 hours, cells were treated with CHx for 0, 2 and 4 hours, and analyzed for fluorescence intensity by FACS Caliber (Becton Dickinson). The fluorescent cells at each time point are represented as a percentage of initial.

FIG. 8

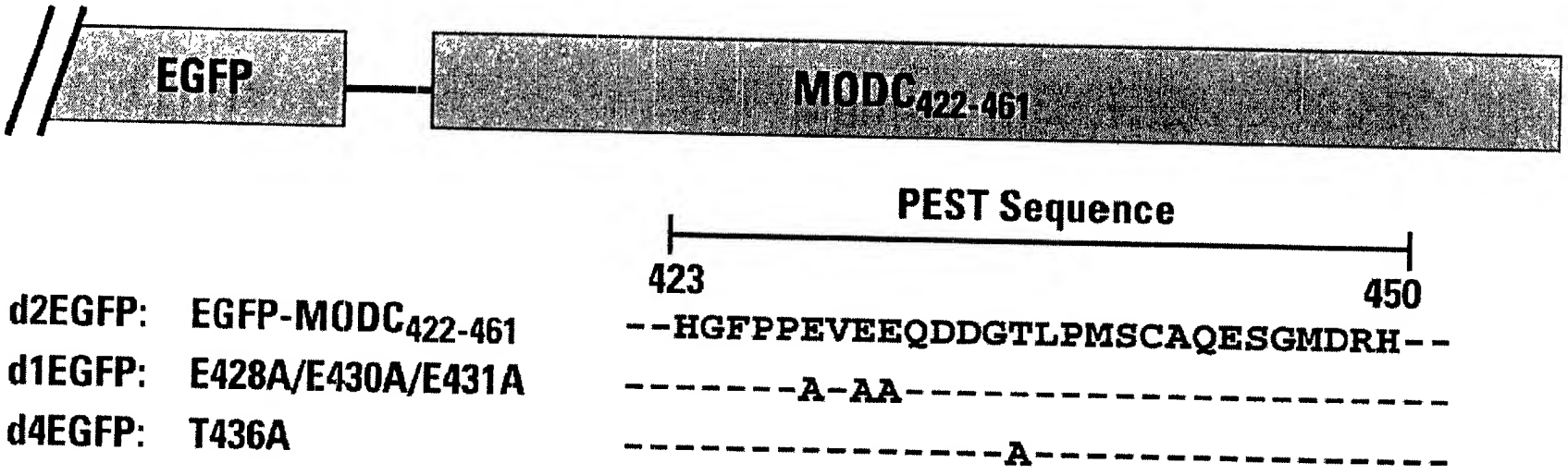
EGFP
MODC422-461
PEST Sequence
423
450
d2EGFP: EGFP-MODC422-461
--HGFPPEVEEQDDGTLPMSCAQESGMDRH--
d1EGFP: E428A/E430A/E431A
-------A-AA--------------------
d4EGFP: T436A
--------------A---------------

FIG. 9

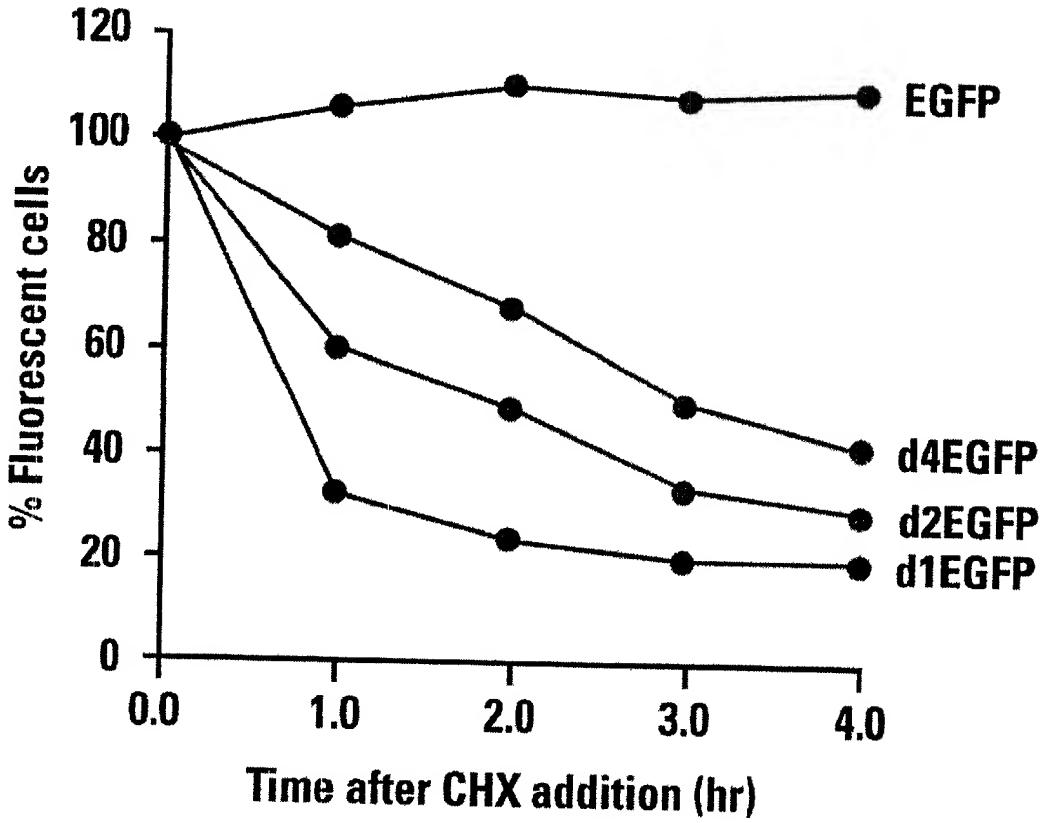

FIG. 10

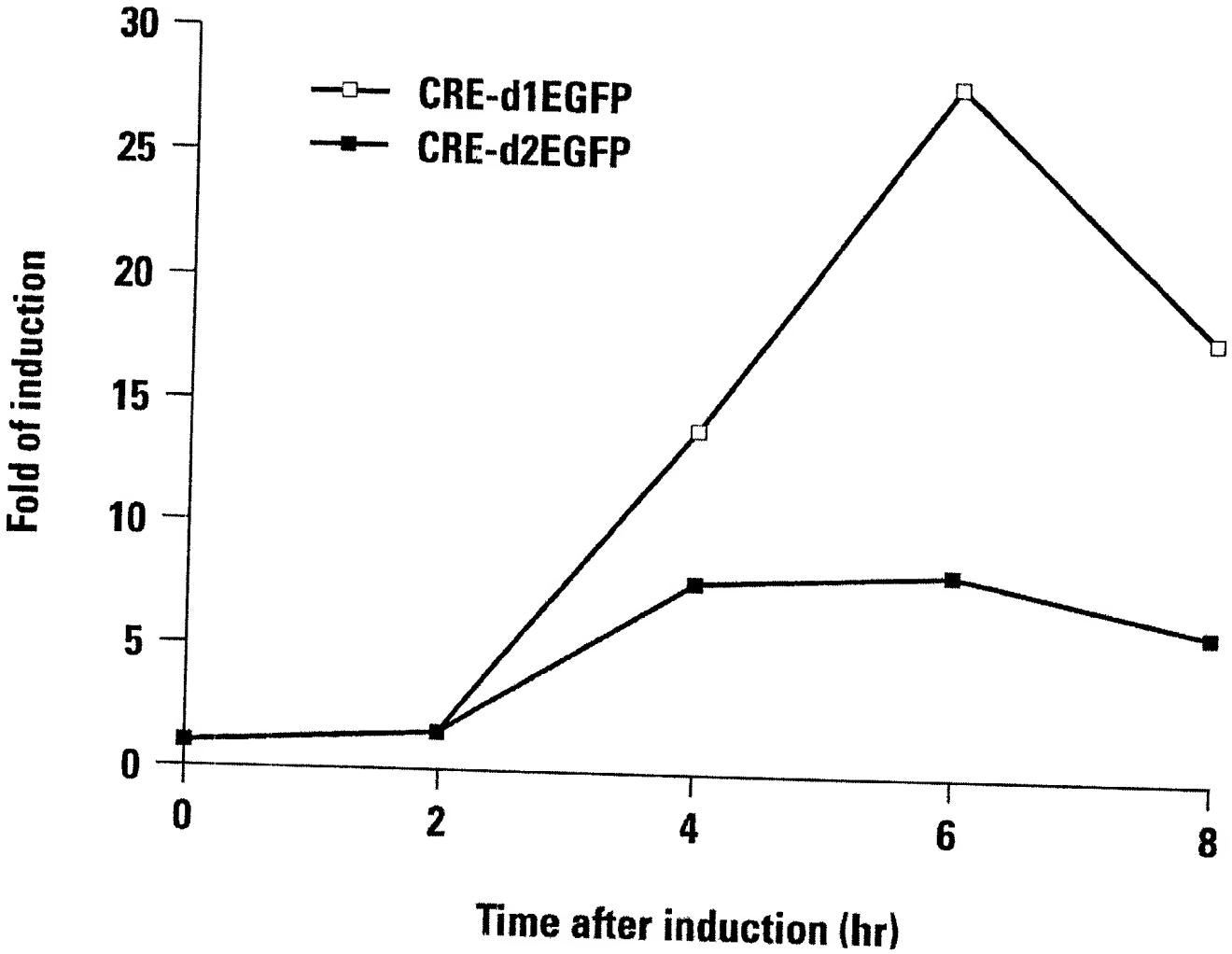

FIG. 11